

FIG. 1

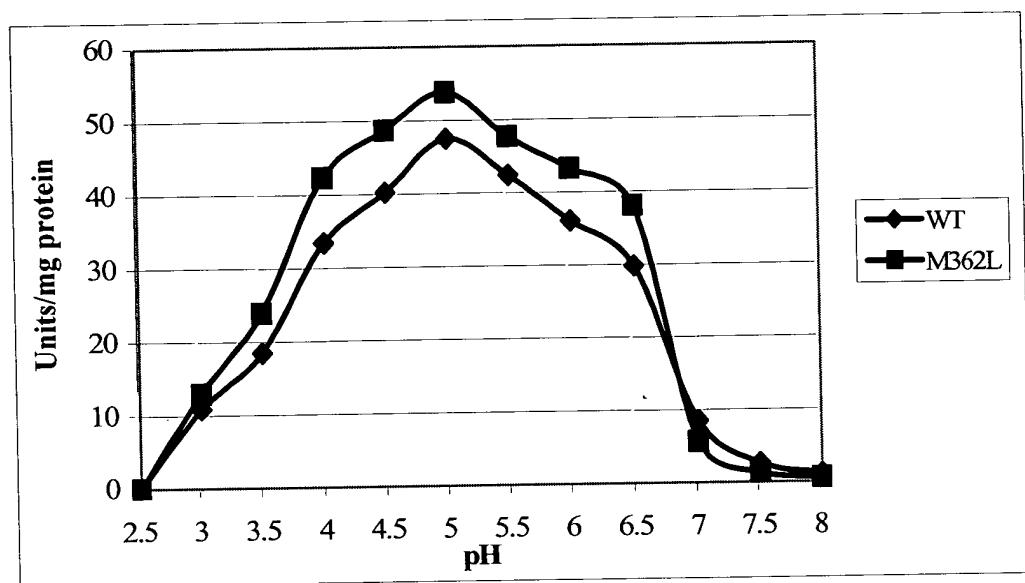


FIG. 2

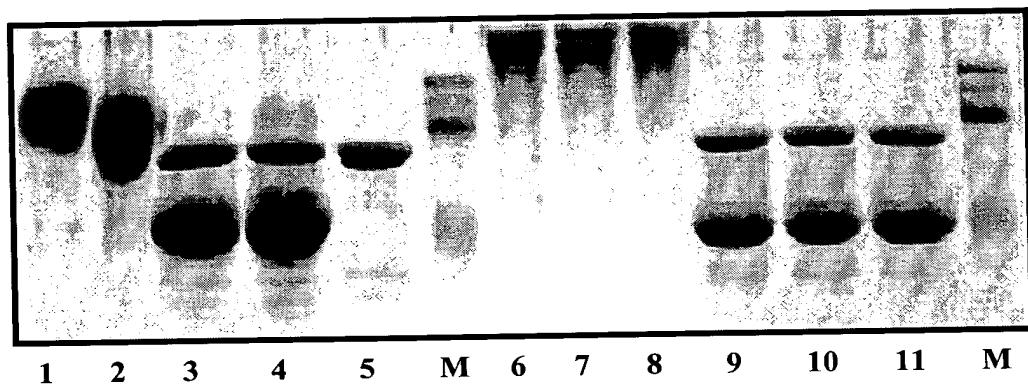


FIG. 3

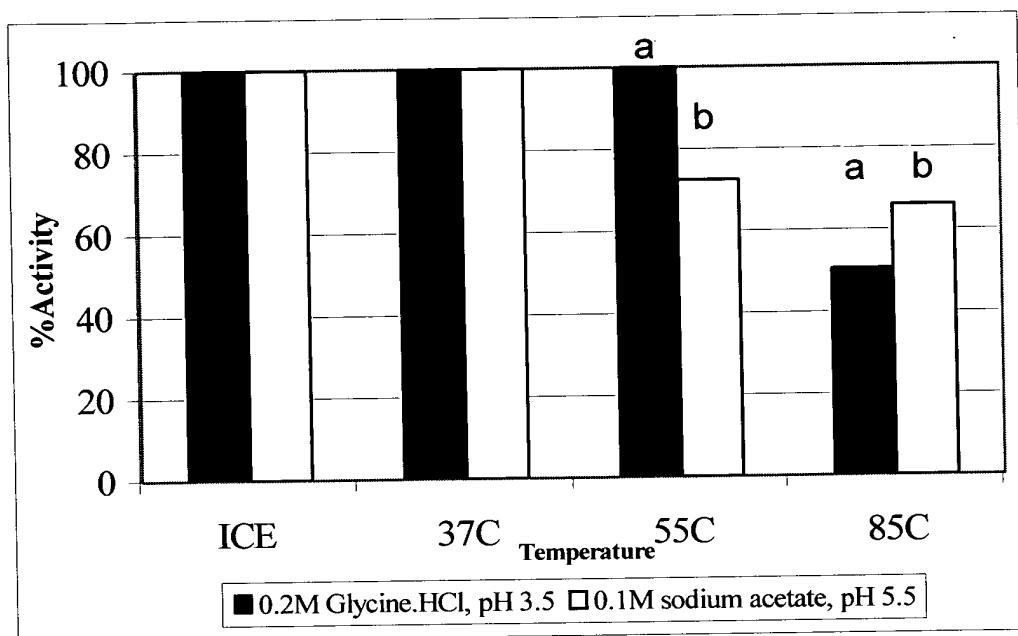


FIG. 4

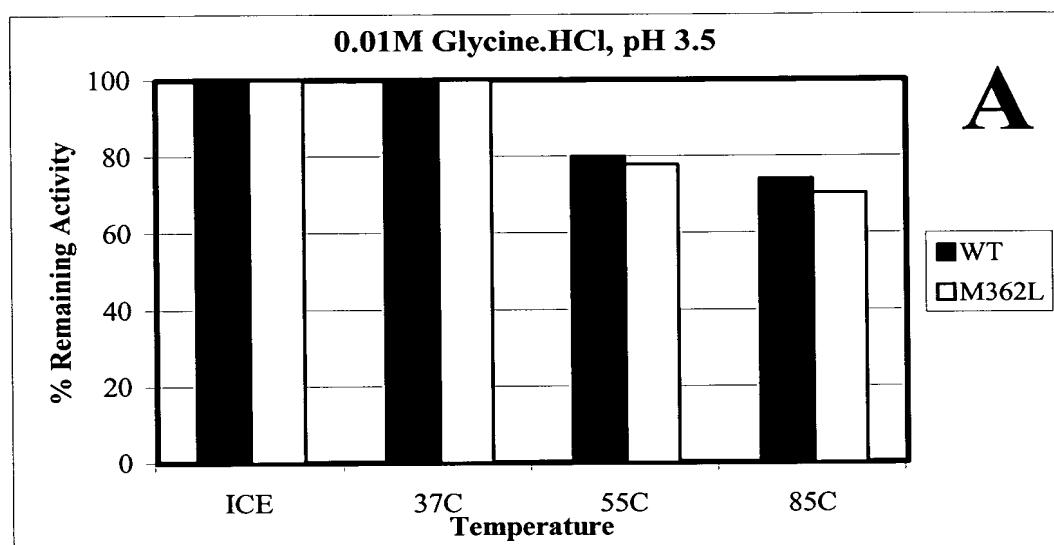


FIG. 5A

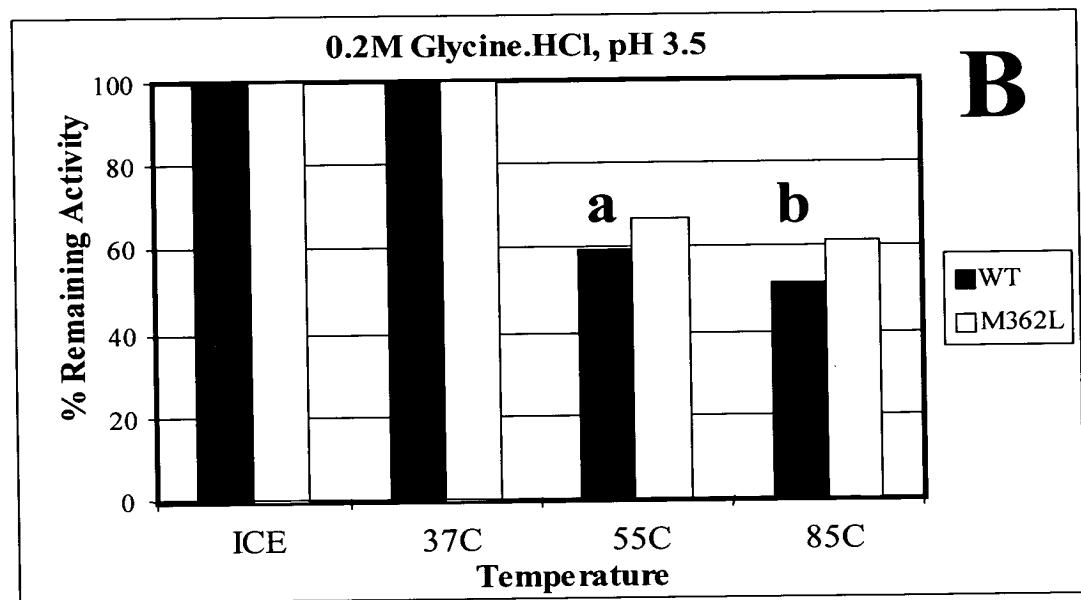


FIG. 5B

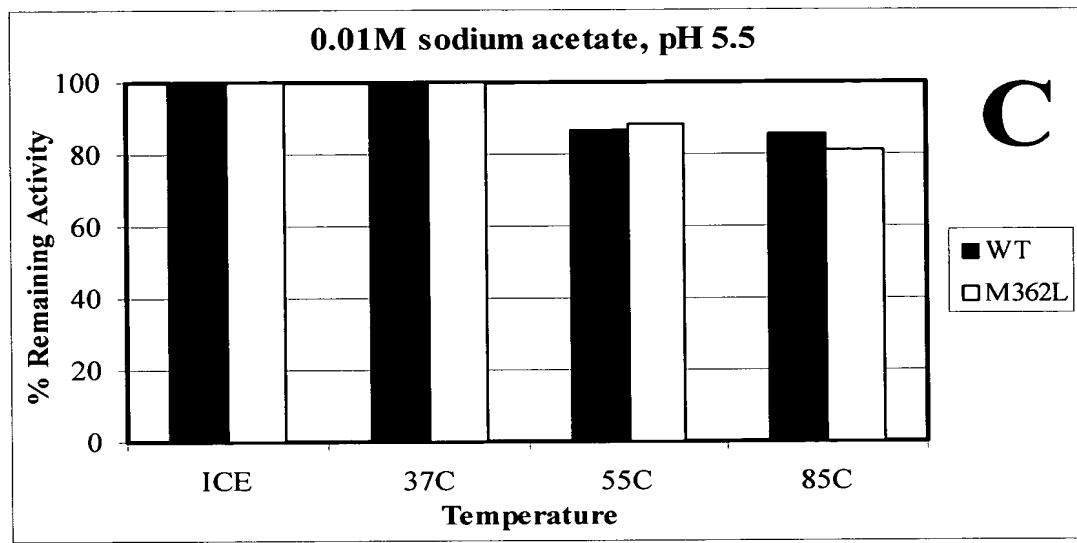


FIG. 5C

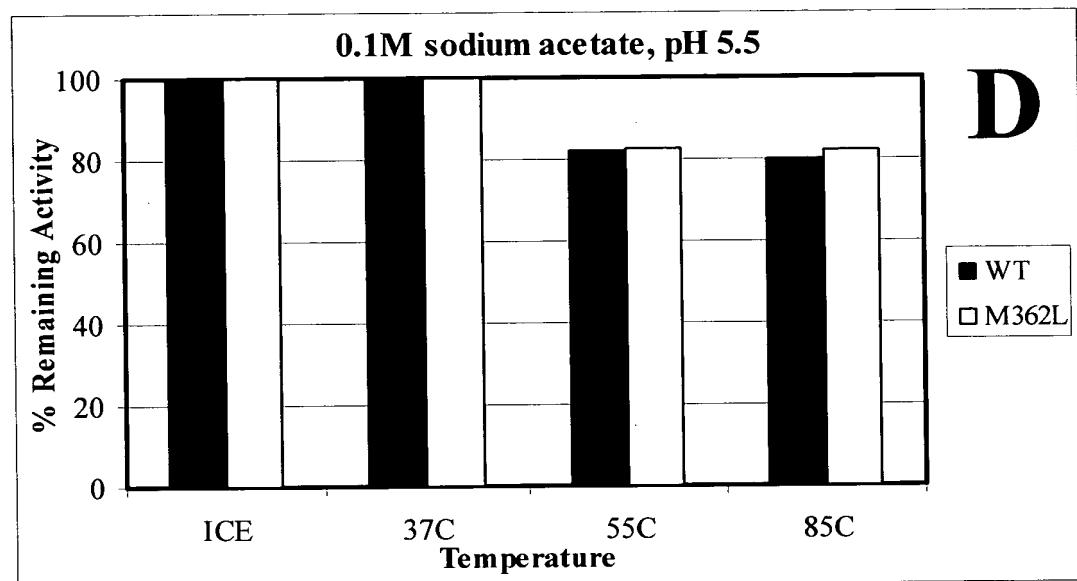


FIG. 5D

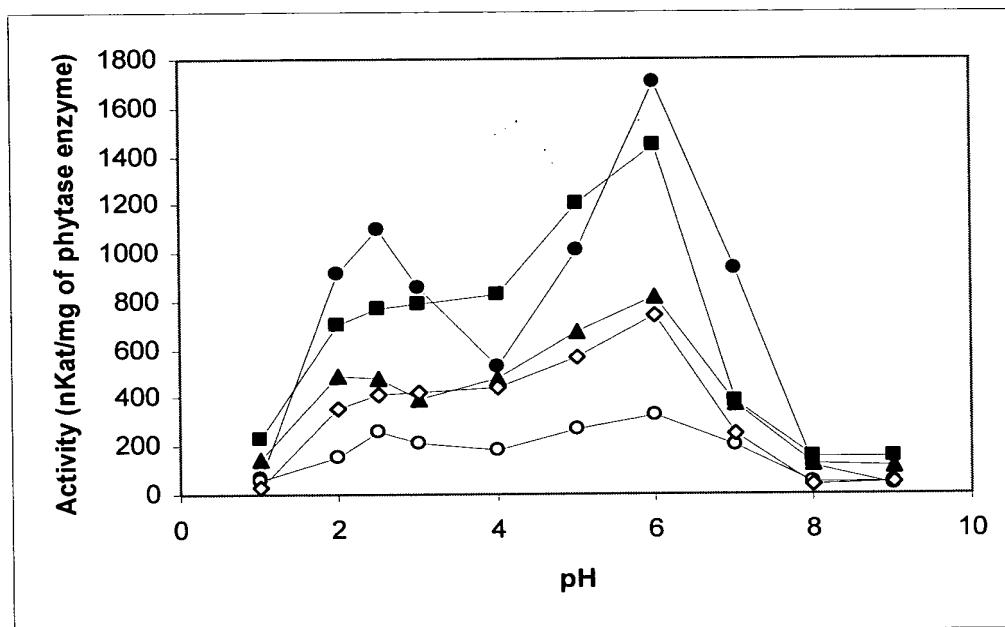


FIG. 6

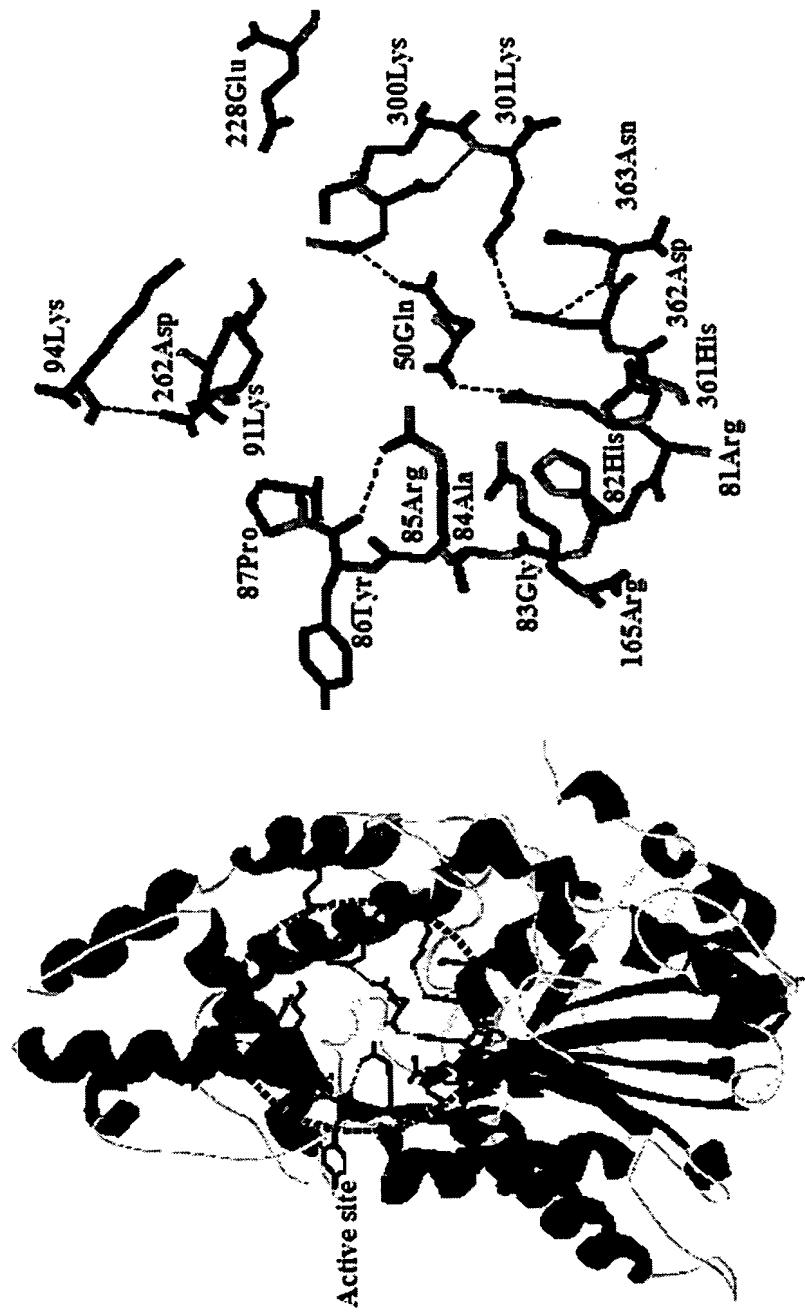


FIG. 7A
FIG. 7B

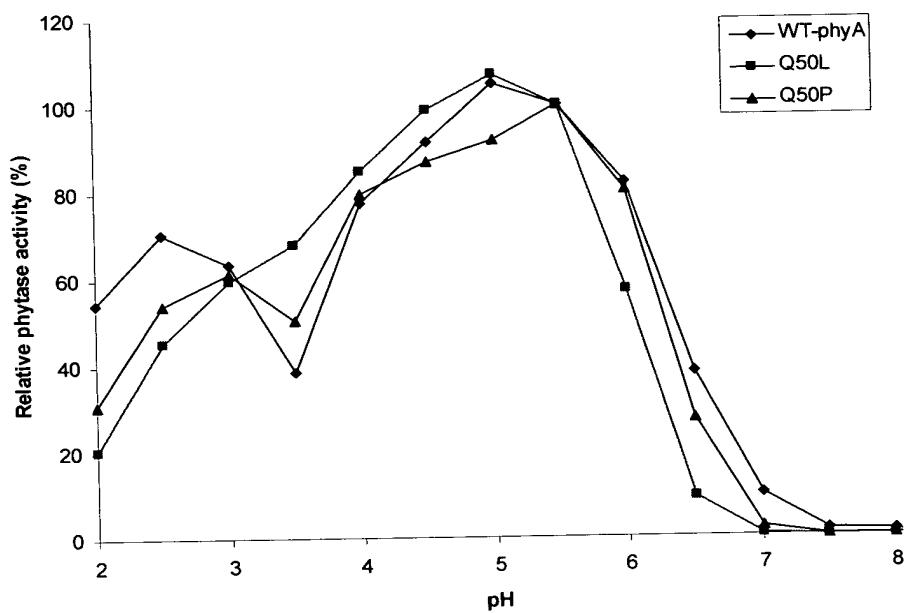


FIG. 8A

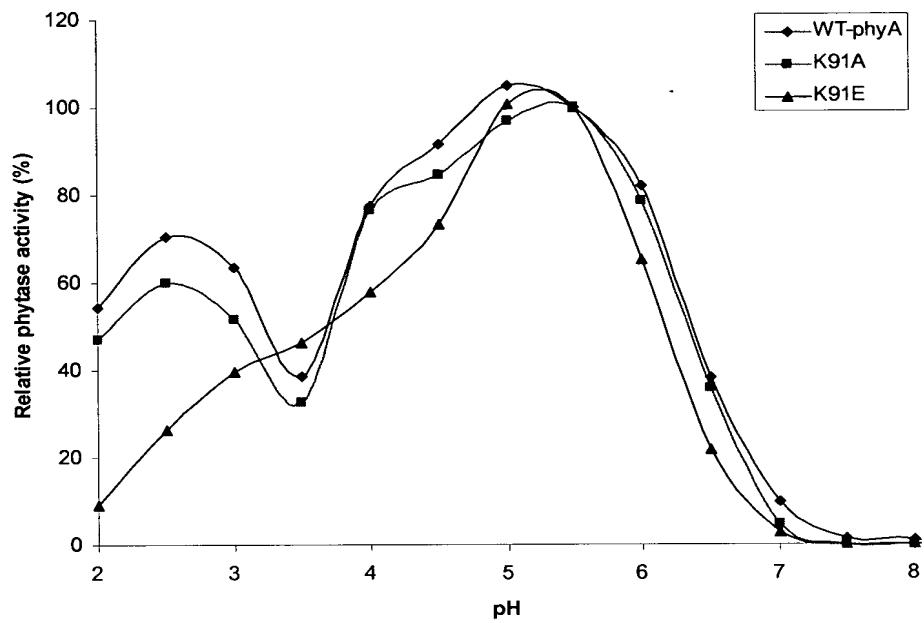


FIG. 8B

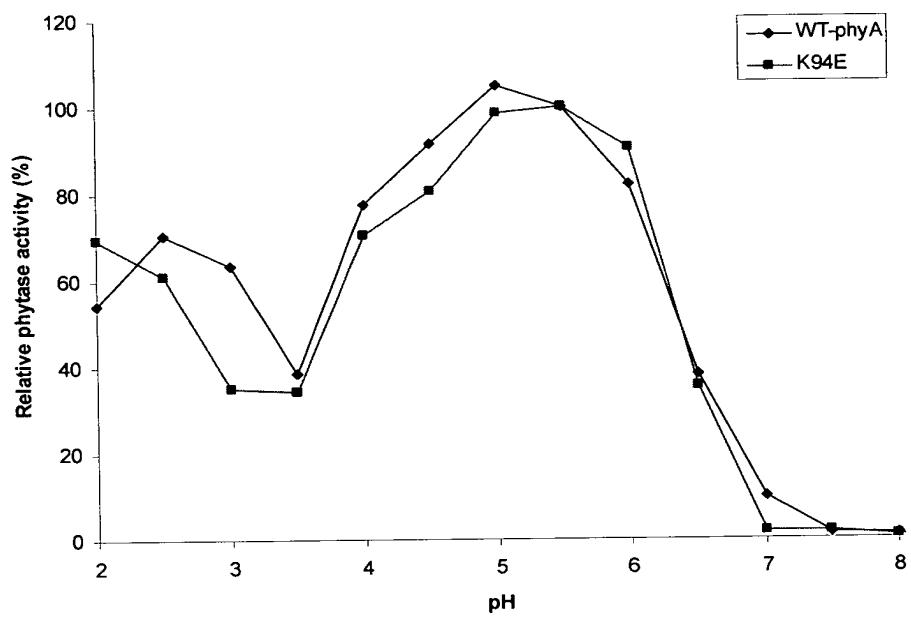


FIG. 8C

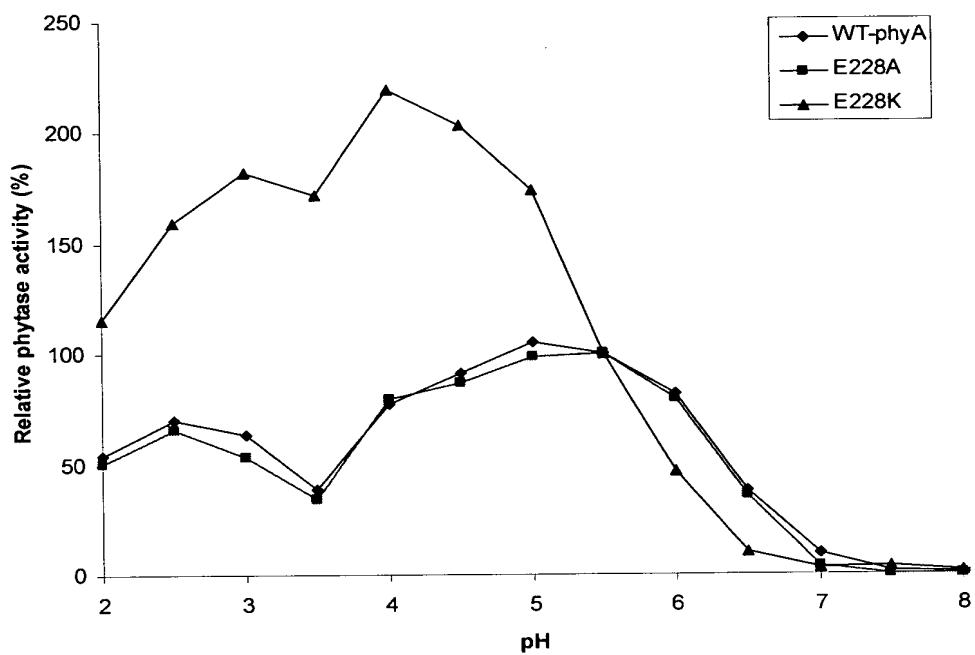


FIG. 8D

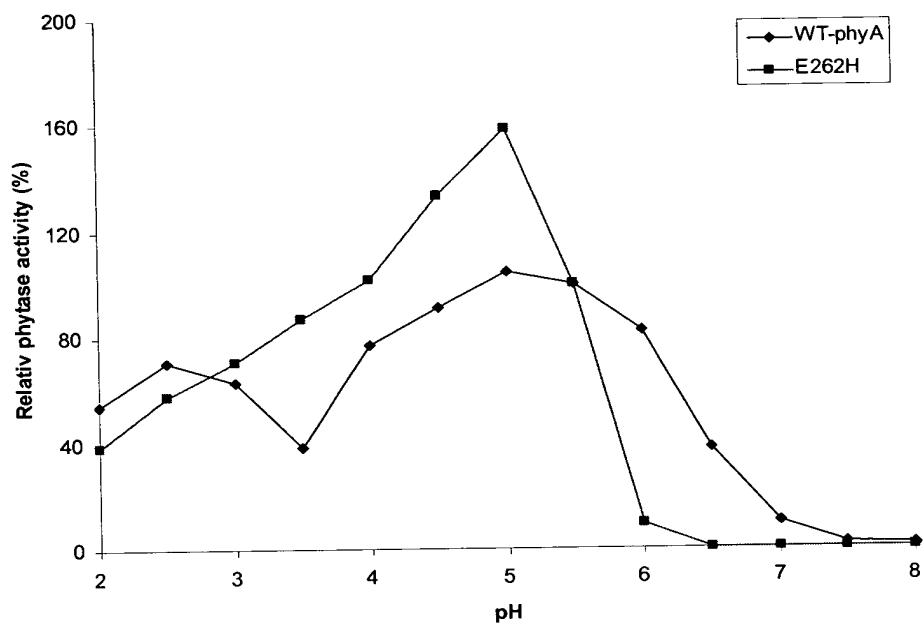


FIG. 8E

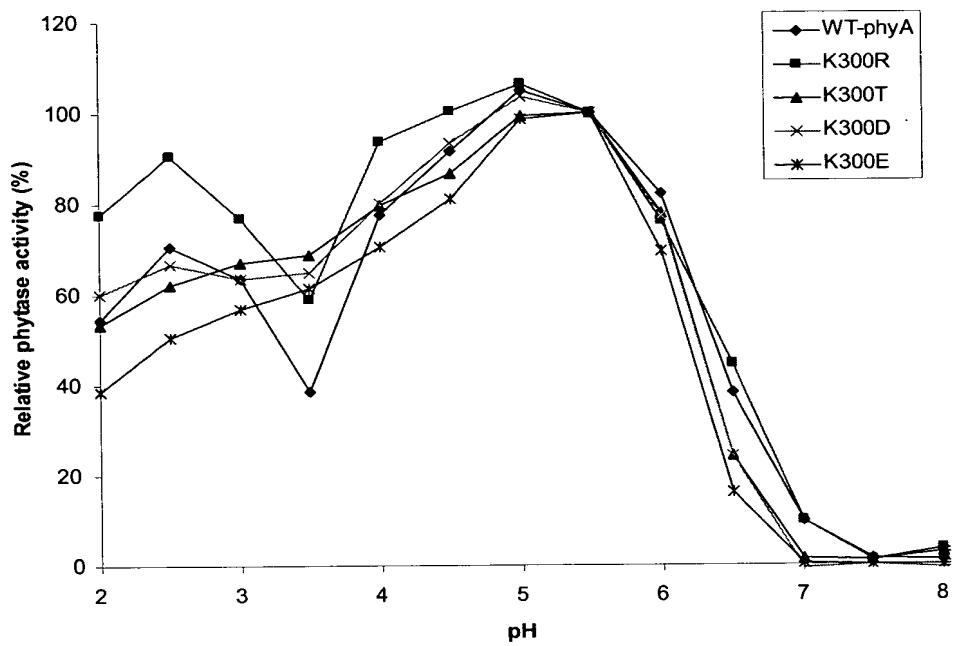


FIG. 8F

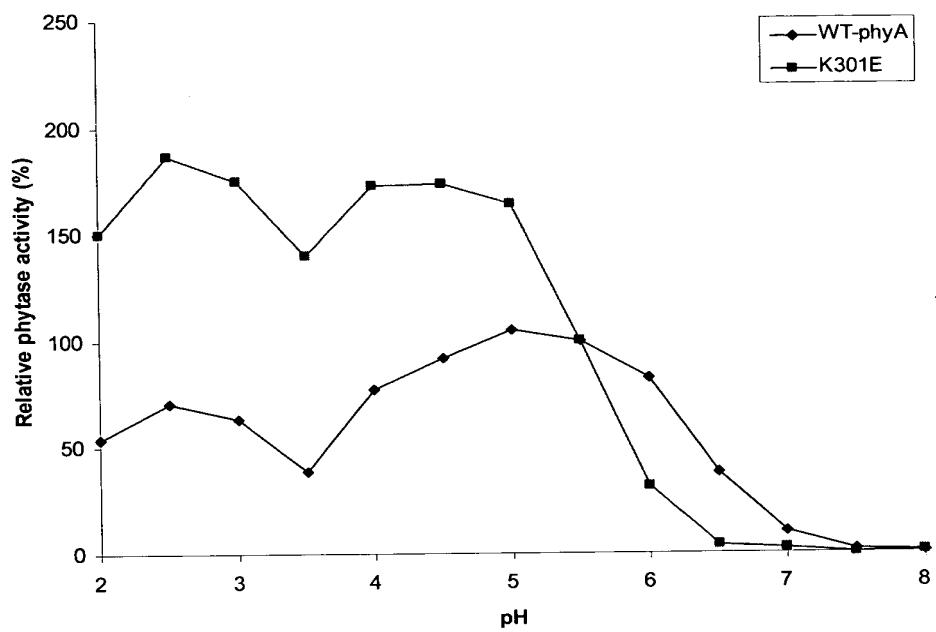


FIG. 8G

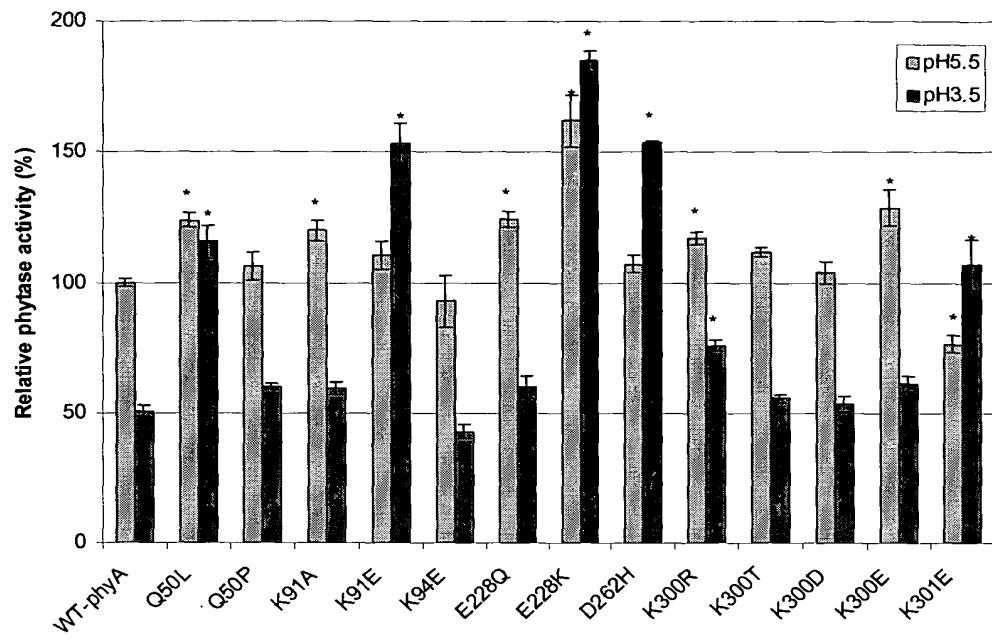


FIG. 9

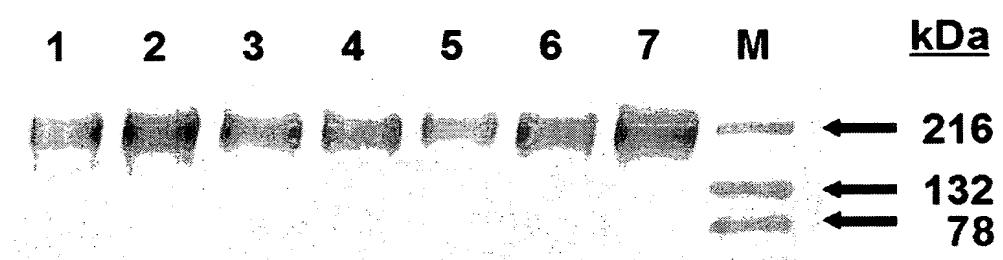


FIG. 10A

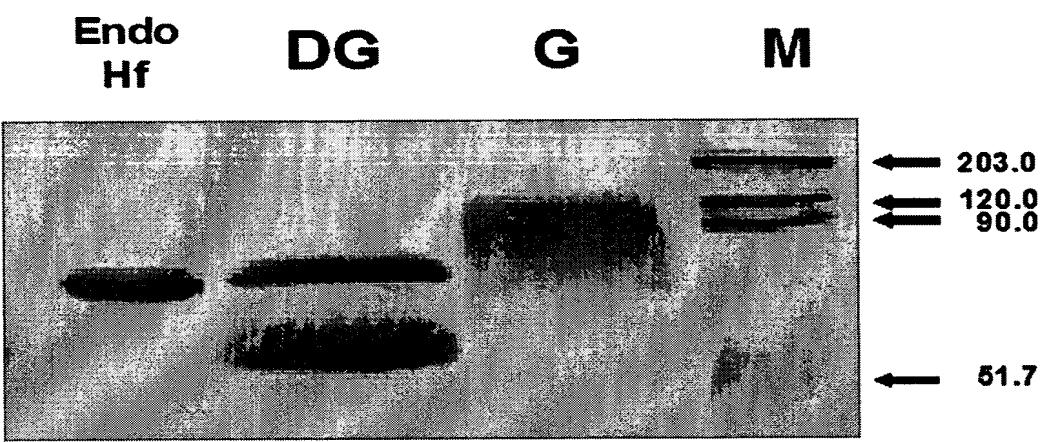


FIG. 10B

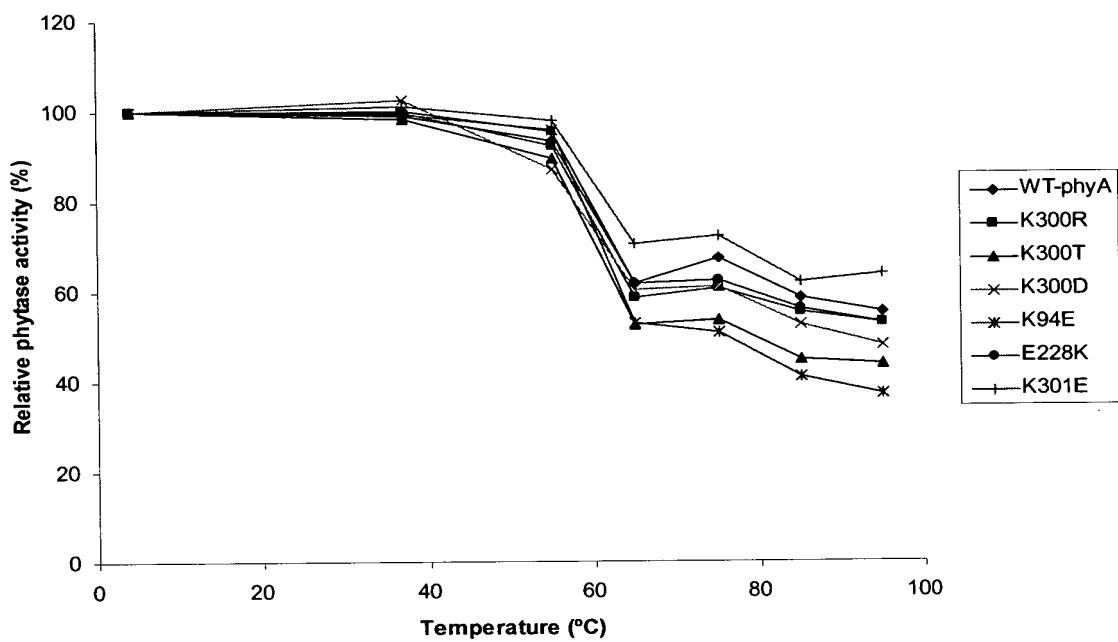


FIG. 11

1st line: Asp. terreus phytase
 2nd line: Asp. niger phytase
 3rd line: Asp. fumigatus phytase

1	MGVFVVLLSI	ATLFGSTS GT	ALGPRGNHSD	CTSVDRGYQC	EPELSHKWGL
1	MGVSAVLLPL	YLLSGVTSGL	AVPASRNQSS	CDTVDQGYQC	FSETSHLWQG
1	MVTLTFLLSA	AYLLSGRVS-	AAPSSAGSKS	CDTVDLGYQC	SPATSHLWQG
****				SSS	SSSS
51	YAPYFSLQDE	SPFPLDVPDD	CHITFVQVIA	RHGARSPTDS	KTKAYAAATIA
51	YAPFFSIAHNE	SVISPEVPAG	CRVTFQAVL S	RHGARYPTDS	KGKKYSAHIE
50	YSPFFSIEDE	LSVSSKLPKD	CRITLVQVLS	RHGARYPTSS	KSKKYKKLVT
****	SSS	S	SSSSSSSSSSSS	SSS	HHHHHHHHHH
101	ATQK NATALP	GKYAFLKSYN	YSMGSENI NP	FGRNQLQDLG	AQFYRRYDTI
101	EI QOONATTED	GKYAFLKTYN	YSLGADDLTP	FGEQELVNSG	IKEYORYESI
100	ATQANATDFK	GKFAFLKTYN	YTLGADDLTP	FGEQQLVNSG	TKFYORYKAI
****	HHHH	HHHH	SSSH	HHHHHHHHHH	HHHH HH
151	TRHINPPEVRA	ADSSRVHESA	EKEVEGEONA	ROGDPHANPH	QPSPERVVV
151	TRNIVPFIRS	SGSSRVIASG	KKFIEGFOST	KLKDPRAQPG	QSSPKIDVV
150	ARSVVPETRA	SGSDRVIASG	EKFIEGFOQA	KLADPGA-TN	RAAPATSVII
****	H	SSS	SS HHHHHHHH	HH	SSS
201	PEGTAYNNTL	EHSIC TAFEA	STVGDAAAADN	FTAVFAPATA	KRLEADLPGV
201	SEASSSNNTL	DPGTCTVFED	SELADTV EAN	FTATEEVESIR	QRLENDLSGV
199	PESETFNNTL	DHGVC TKEA	SQLGDEVAAN	FTALFAPDIR	ARAEKHLPGV
****		HHHH	HHHHHHHH	HHHH	HHHHHH
251	QISADDVVNI	MAMCPETV S	LTD DAHTLSP	FCDLFTAAEW	TOYNVILSLD
251	LTIDTEVTVL	MDMCSFDTIS	TSTVDTKLSP	FCDLFTHDEW	INYDYLQSLK
249	LTIDDEVVSL	MDMCSFDTVA	RTSDASQLSP	FCOLFTHNEW	KKYNYLQSLG
****	HHHHHH	HHHHHHH	HH	HHHH	HHHHHHHHHH
301	KYYGYGGGNP	LGPVQGVGWA	NELIARLTRS	PVHDHTCVNN	TLDANPATFP
301	KYYGHAGNP	LGPTQGVGYA	NELIARLTHS	PVHDDTSSNH	TLDSSPATFP
299	KYYGYGAGNP	LGPAQGIGFT	NELIARLTRS	PVQDHTSTNS	TLVSNPATFP
****	H	HH	HHHHHHHH	H	HHH
351	LNATLYADFS	HDSNLVSI FW	ALGLYNGTKP	LSOTTVEDIT	RTDGYAAAWT
351	LNSTLYADFS	HDNGIISILF	ALGLYNGTKP	LSTT TVENIT	QTDGFSSAWT
349	LNATMYVDF	HDNSMVSTFF	ALGLYNGTEP	LSRTSVE SAK	ELDGYSASW
****	SSSSSS	HHHHHHHH	H		
401	VPFAARAYIE	MMOCRAEKOP	LVRV L VNDRV	MPLHGC AVDN	LGRCKRD DFV
401	VPFASRLYVE	MMOCQAE OFP	LVRV L VNDRV	VPLHGC PVD	LGRCTR DSEV
399	VPFGARAYFE	TMOCKSEKEP	LVRALINDRV	VPLHGC DVDK	LGRCKLND FV
****	SSSSSS	SSSS	S	SSSSSS SS	SSHHHH
451	EGLSFARAGG	NWAECF			
451	RGLSFARSGG	DWAECFA			
449	KGLSWARSGG	NWGECEFS			
****	H	HHH	HHGTT		

*Red letter shows the mutation site for substrate binding site.
 *Bold letters are known as critical catalytic active sites.

FIG. 12

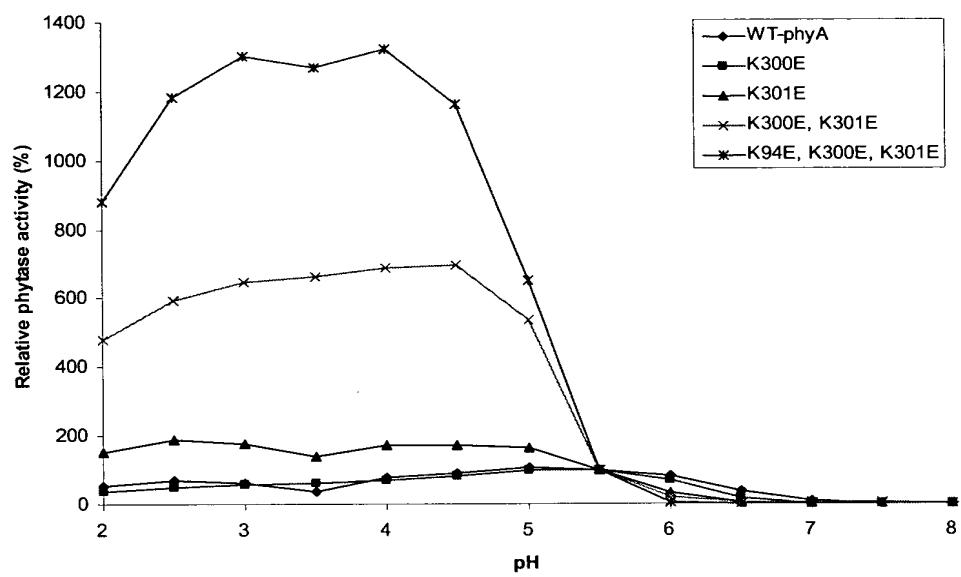


FIG. 13A

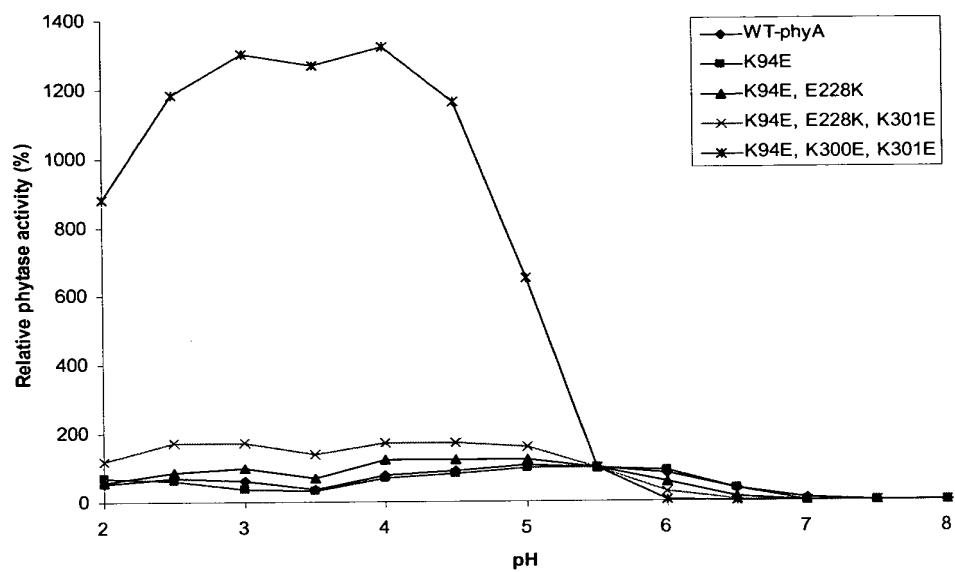


FIG. 13B

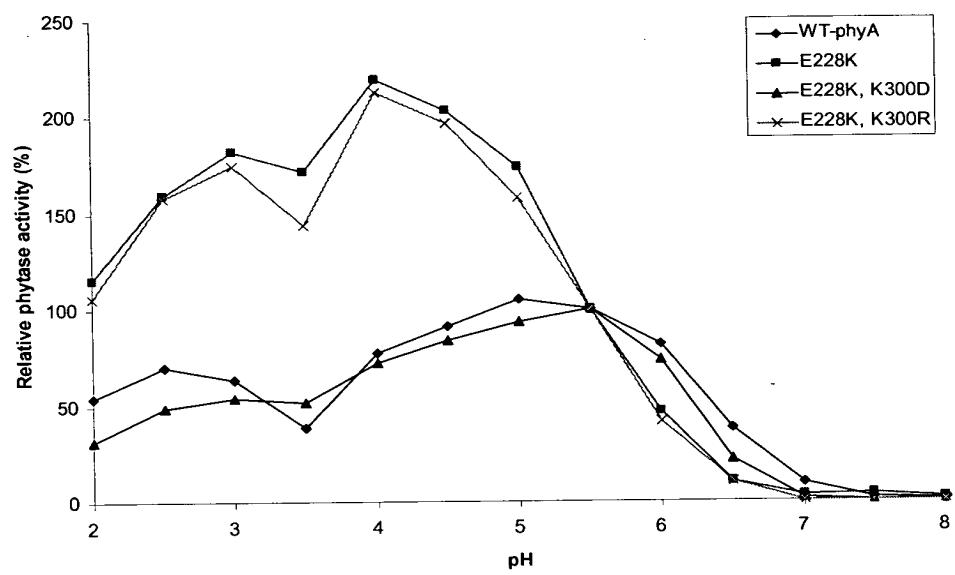


FIG. 13C

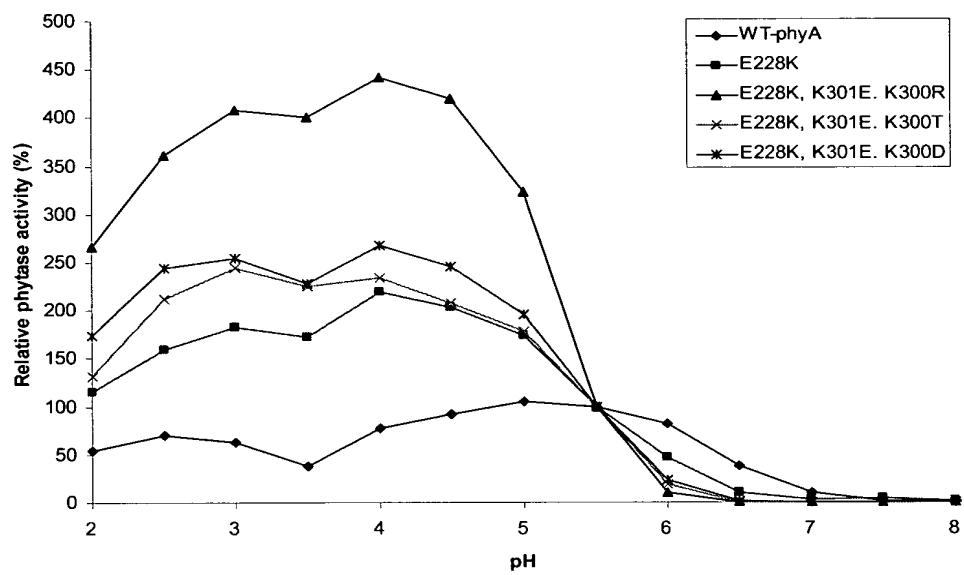


FIG. 13D

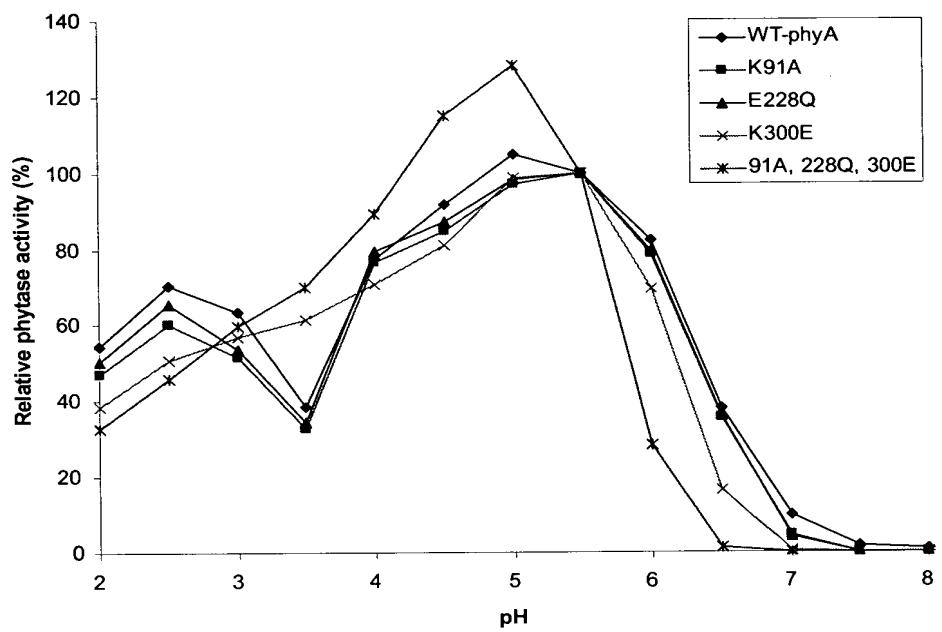


FIG. 13E

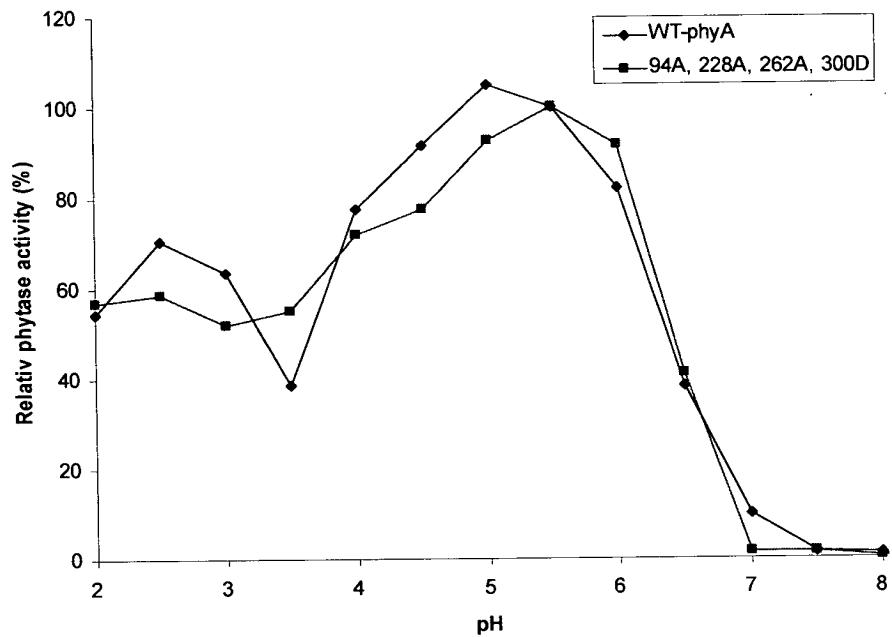


FIG. 13F

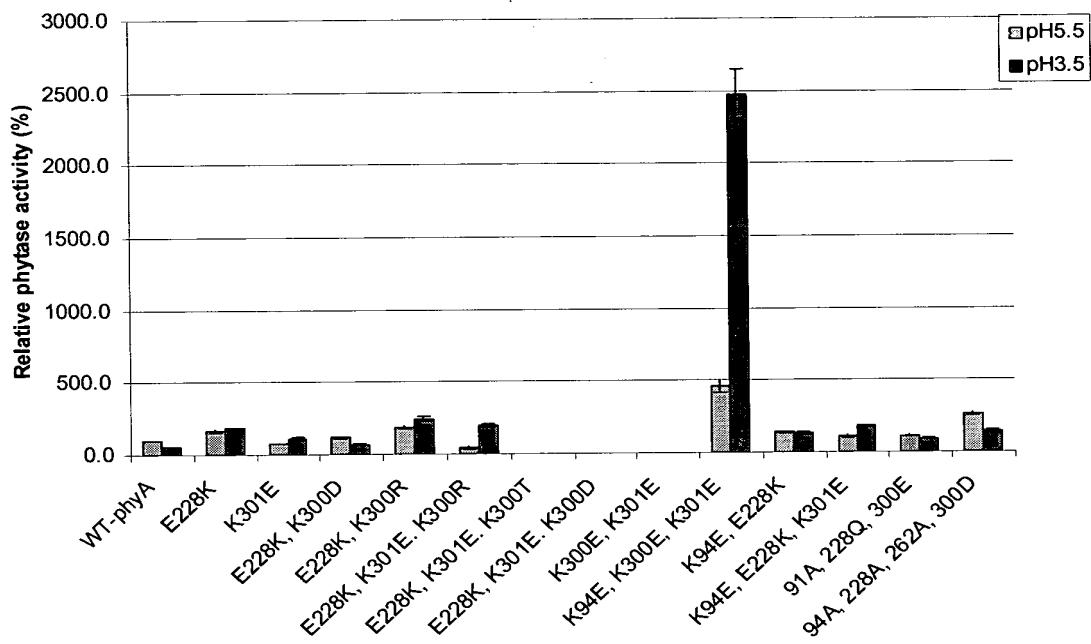


FIG. 14A

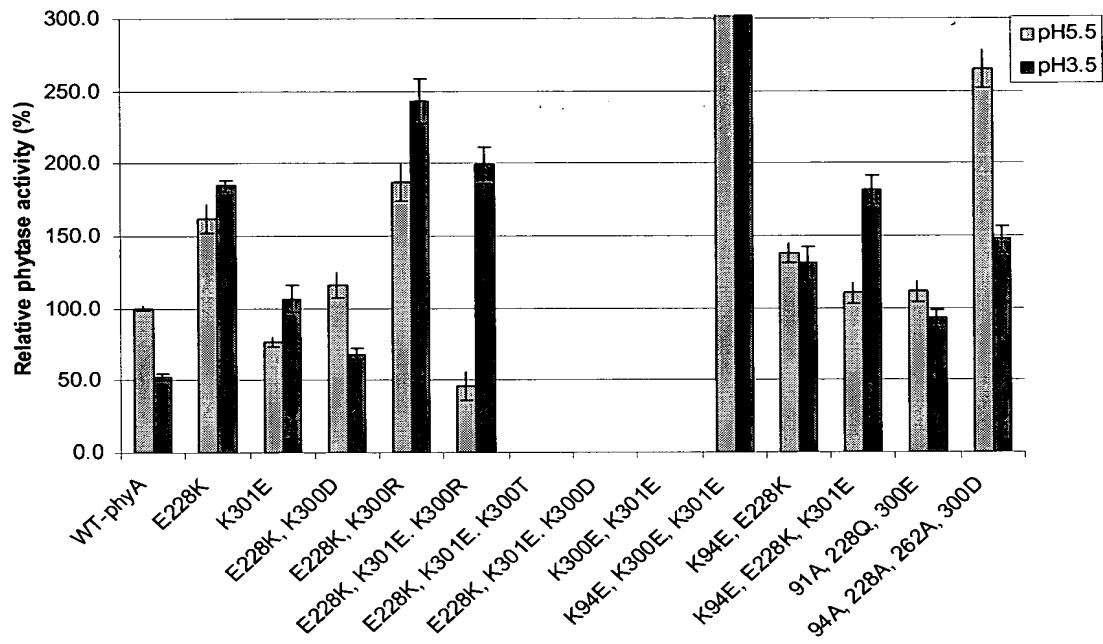


FIG. 14B

Plasma Inorganic Phosphate (PIP) of Pigs Fed Low-P Diets

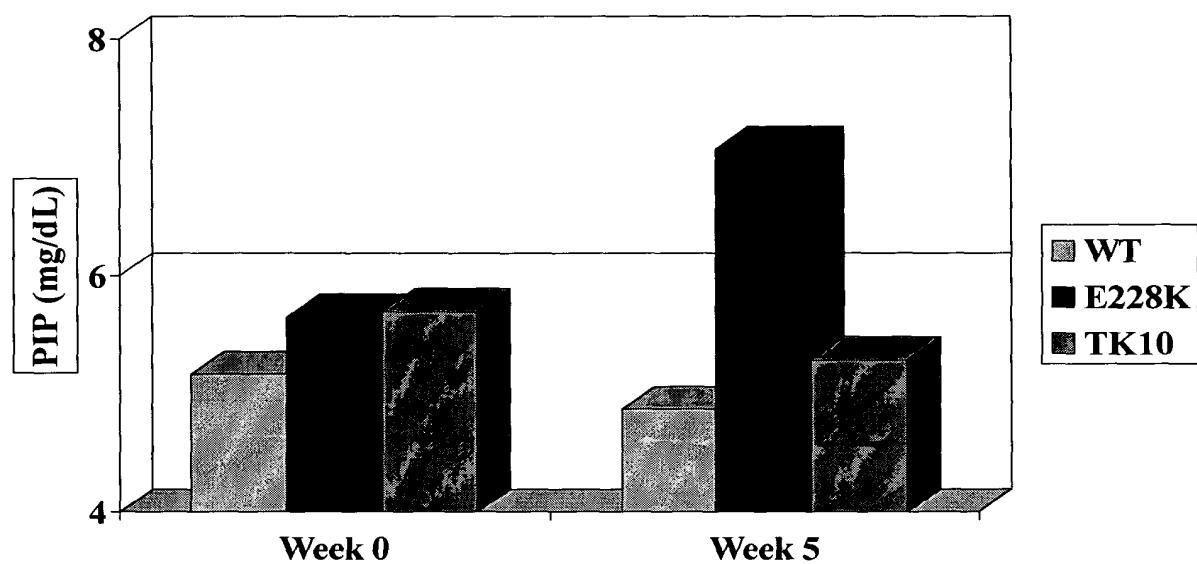


FIG. 15

Plasma Alkaline Phosphatase (AKP) Activity of Pigs Fed Low-P Diets

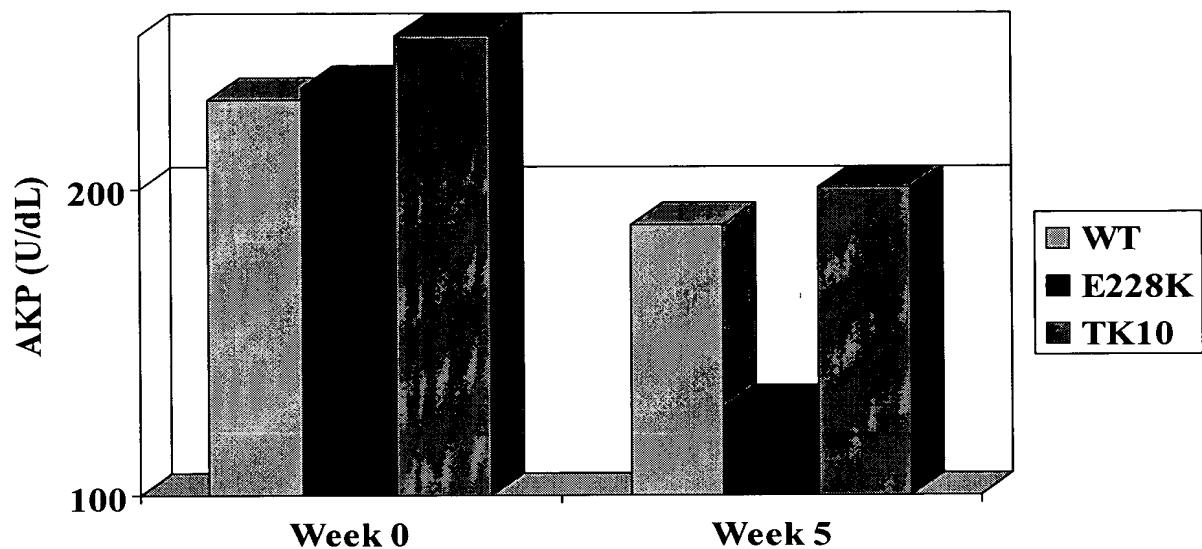


FIG. 16

Average Daily Gain (ADG) of Pigs Fed Low-P Diets

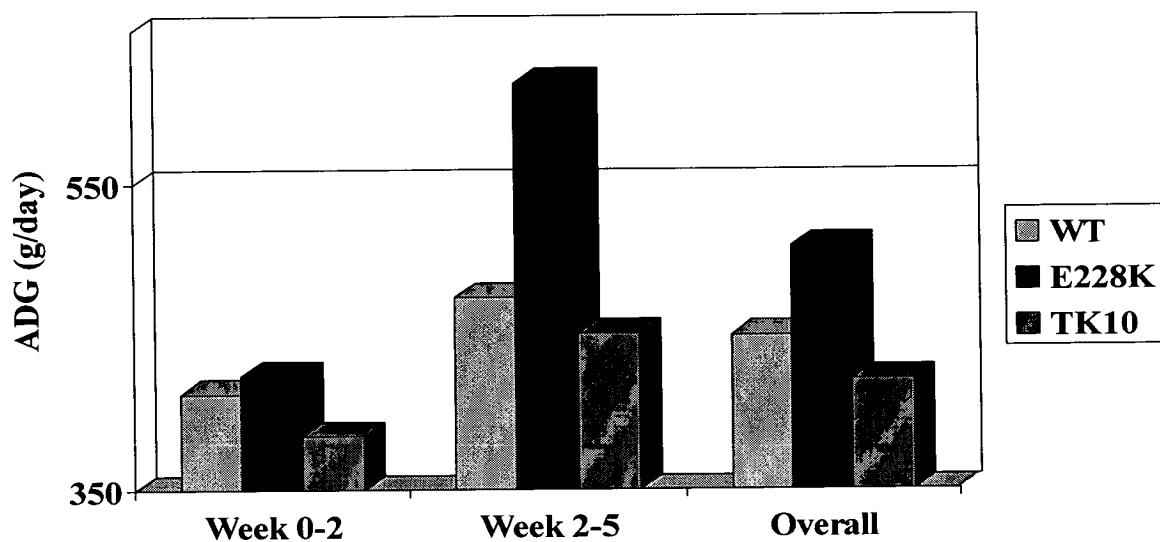


FIG. 17

Gain/Feed of Pigs Fed Low-P Diets

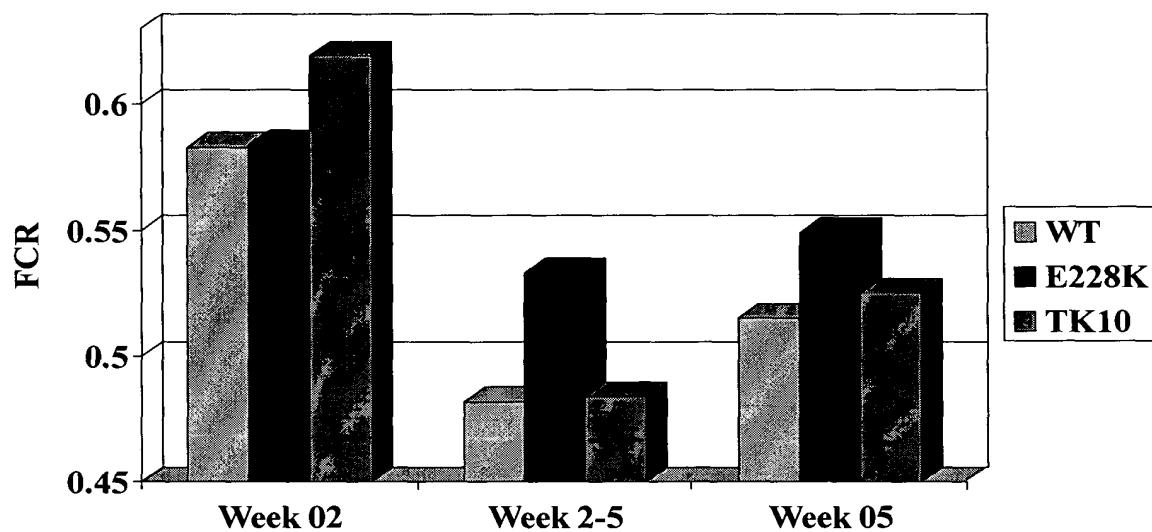


FIG. 18